



#4

SEQUENCE LISTING

<110> Ben-Sasson, Shmuel

<120> Tissue Remodeling

<130> BEN-SASSON=7

<140> US 10/032,330

<141> 2001-12-31

<150> PCT/US00/32852

<151> 2000-12-04

<150> US 09/161,094

<151> 1998-09-25

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<170> PatentIn version 3.1

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Gln Thr Leu

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Xaa Gly Glu Ser Val Pro Glu Tyr Gln Met Ala Phe
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Glu Ile Ala Arg Arg Cys Ser Ile Gly Gly Ile His Glu Asp Tyr Gln
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Leu Pro Tyr Tyr Asp Leu Val Pro Ser Asp Pro Ser Val Glu Glu

20

25

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<400> 33

Glu	Leu	Val	Ser	Arg	Cys	Lys	Ala	Ala	Asp	Gly	Pro	Val	Asp	Glu	Tyr
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Met	Leu	Pro	Phe	Glu	Glu	Glu	Ile	Gly	Gln	His	Pro	Ser	Leu	Glu	Glu
			20					25					30		

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Glu	Ile	Phe	Met	Arg	Cys	Thr	Asp	Leu	Phe	Pro	Gly	Glu	Ser	Val	Pro
1				5				10						15	
Glu	Tyr	Gln	Met	Ala	Phe	Gln	Thr	Glu	Val	Gly	Asn	His	Pro	Thr	Phe
			20					25					30		

Glu Asp

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Glu	Met	Thr	Ser	Arg	Cys	Asn	Ala	Val	Gly	Glu	Val	Lys	Asp	Tyr	Glu
1				5					10					15	
Pro	Pro	Phe	Gly	Ser	Lys	Val	Arg	Glu	His	Pro	Cys	Val	Glu	Ser	
			20					25					30		

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<400> 36

Glu Ile Ala Arg Arg Cys Asn Ser Gly Gly Val His Glu Glu Tyr Gln
1 5 10 15

Leu Pro Tyr Tyr Asp Leu Val Pro Ser Asp Pro Ser Ile Glu Glu
20 25 30

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<400> 37

Glu Met Ala Arg Arg Cys Ile Thr Gly Gly Ile Val Glu Glu Tyr Gln
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Leu Pro Tyr Tyr Asn Met Val Pro Ser Asp Pro Ser Tyr Glu Asp
20 25 30

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Glu Ile Ala Arg Arg Thr Ile Val Asn Gly Ile Val Glu Asp Tyr Arg
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Pro Pro Phe Tyr Asp Val Val Pro Asn Asp Pro Ser Phe Glu Asp
20 25 30

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<400> 39

Ser Asp Tyr His Glu His Gly Ser Leu Phe Asp Tyr Leu Asn Arg Tyr
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Thr Val Thr Val Glu
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Thr Ala Phe His Asp Lys Gly Ser Leu Thr Asp Tyr Leu Lys Gly Asn
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Ile Ile Thr Trp Asn Glu
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Met Glu Tyr Tyr Pro Asn Gly Ser Leu Cys Lys Tyr Leu Ser Leu His
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Thr Ser Asp Trp Val
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<400> 42

Thr Ala Phe His Ala Lys Gly Asn Leu Gln Glu Tyr Leu Thr Arg His
1 5 10 15

Val Ile Ser Trp Glu
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<400> 43

Ser Asp Tyr His Glu His Gly Ser Leu Phe Asp Tyr Leu Asn Arg Tyr
1 5 10 15
Thr Val Thr Ile Glu
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<400> 44

Thr Asp Tyr His Glu Asn Gly Ser Leu Tyr Asp Phe Leu Lys Cys Ala
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Thr Leu Asp Thr Arg
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Thr His Tyr His Glu His Gly Ser Leu Tyr Asp Phe Leu Gln Arg Gln
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Thr Leu Glu Pro His
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<400> 46

Ala Asp Asn Lys Asp Asn Gly Thr Trp Thr Gln Leu Trp Leu Val Ser
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Asp

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<400> 47

Ala Glu Lys Arg Gly Ser Ser Leu Glu Ala Glu Leu Trp Leu Ile Thr
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Ala

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<400> 48

Gly Asp Glu Arg Val Thr Ala Asp Gly Arg Met Glu Tyr Leu Leu Val
1 5 10 15

Met Glu

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Ala Glu Glu Arg Lys Thr Glu Leu Gly Lys Gln Tyr Trp Leu Ile Thr
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Ala

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Asp

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Ser	Asp	Met	Thr	Ser	Arg	Asn	Ser	Ser	Thr	Gln	Leu	Trp	Leu	Ile	Thr
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His

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Ser Thr Pro Gly Met Lys His Glu Asn Leu Leu Gln Phe Ile Ala
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Arg Val Pro Leu Met Glu His Asp Asn Ile Ala Arg Phe Ile Val
1 5 10 15

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<210> 58

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<400> 58

Gln Thr Val Leu Met Arg His Glu Asn Ile Leu Gly Phe Ile Ala
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<400> 59

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